

APPROVED	O.G. FIG.	
	CLASS/SUBCLASS	
BY		
DRAFTSMAN		

FORWARD SEQUENCING PRIMER:

5'-CACTCACGCAAACGG-linker-TGTAAAACGACGGCCAGT-3'

⏟

Capture complement

⏟

Sequencing primer

REVERSE SEQUENCING PRIMER:

5'-AACTCTCCCAAGAGCAC-linker-CAGGAAACAGCTATGACC-3'

⏟

Capture complement

⏟

Sequencing primer

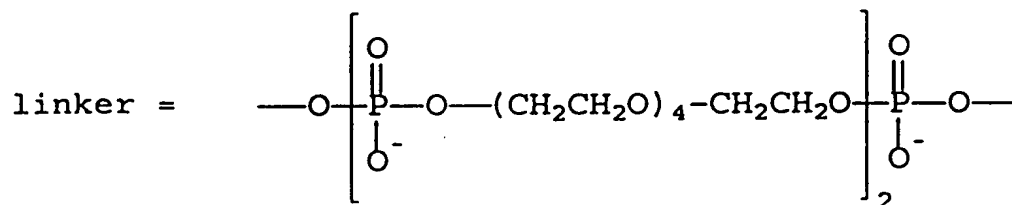


FIG. 1

APPROVED	O.G. FIG.	
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CGAGGAAGAATGAAATGTCGCA GTTGA GTGTGTT CCTCGCATCC CAGGAGCAATGCA GTGCTGATTGTC TGTTCGTGAGATA CCCTTTTGT
450 460 470 480 490 500 510 520 530

TCGGAAACATA AGGTTTAAAGAA ACACACACAGTTT TCTGGGGTC TCTTGTTAGATGTAA ATGTCCGCCA TCCA GACCC TTACTA TCGA GTGTA
540 550 560 570 580 590 600 610 620

ACTCAATTG GGTTACTAATCACCCCG CCCACCCGCAACAA TCA GGA TCA TTGGA ATGGGA ACCTCA TTCAA CT TCTAGGCTGAGAGTA
630 640 650 660 670 680 690 700 710

TTGCCTTCA ACCTA TTTCA TTTCTAGACTAG CACCA TTCAACT GAGA TAA TAAAGCAACAA TAA CCNIT GATCA TGA TGIC TA CCAA AAA
720 730 740 750 760 770 780 790 800

ATGIGCCCCGGAAAAGTCTTTTA GGAATTGTAA TGAGAAAANA AA TGAAAATGGGGCA TAAATTC TCTT
810 820 830 840 850 860 870

FIG. 2A2

APPROVED	O.G. FIG.	
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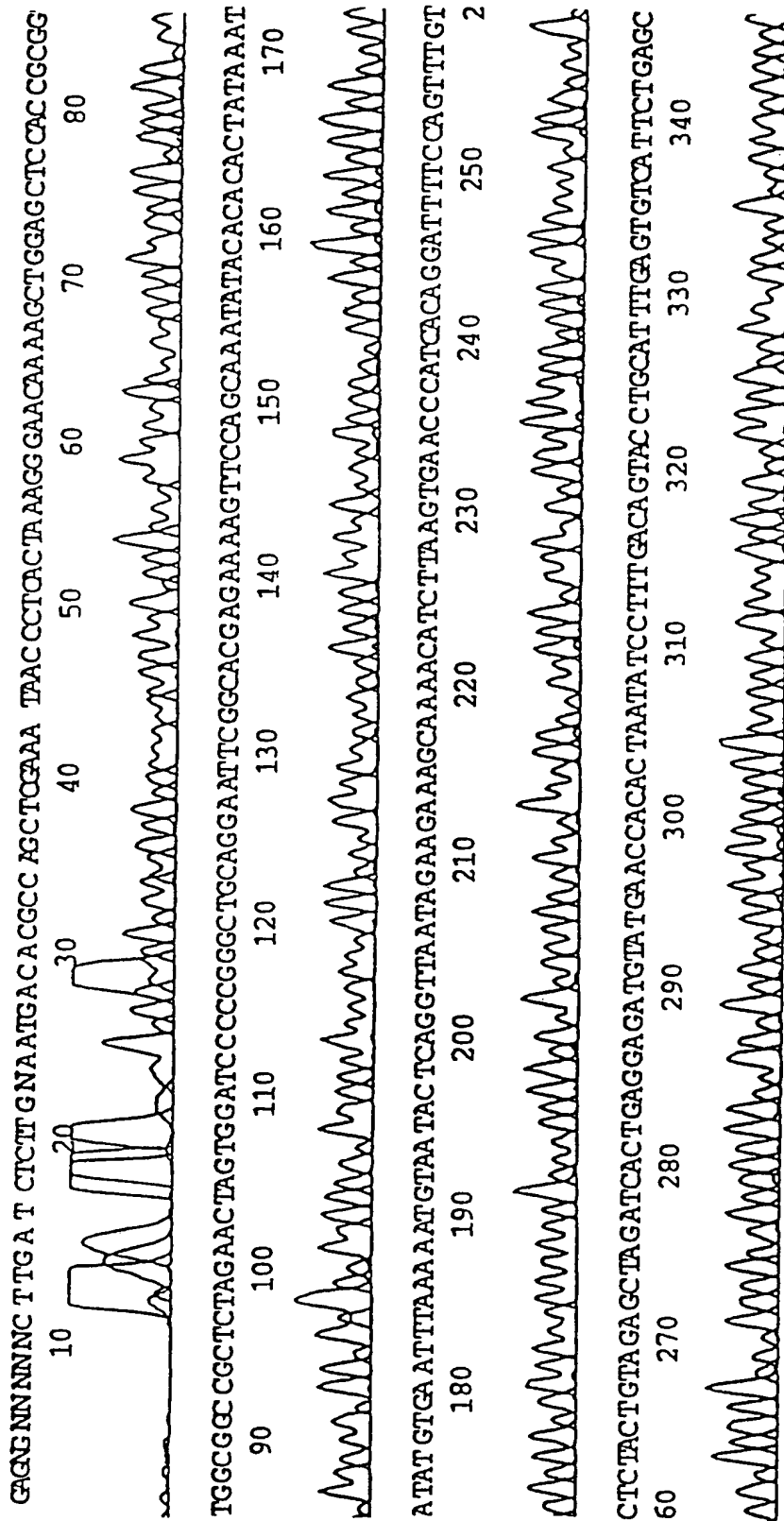


FIG. 2B1

APPROVED	O.G. FIG.	
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AAATTGAGTAA TTTTACTCAATGAGGTCATCATCTTCAC TAGCTTTTGIGGTT TTTTCTTTCTTT TTTTCTTTT TTTTITTAGG G
 350 360 370 380 390 400 410 420 430



GAAAGGGTGGTACA TAAAGGAC TTCTTATGTAAC TTTATTGAAGTCAAATCATGAC CCTTACTGTGCAATTTTATG GAAGCATTTAAAAT
 440 450 460 470 480 490 500 510 520



TATCTTGTCAC TAATTGCA TTA TTGTAA GAAC TACCTCAGATTTTGTGA GTAATATGAAGAAAAGTC TATGTTCA TTAACAGCTACTA
 530 540 550 560 570 580 590 600 610



AGTTCTAGANGTGTTTT GAAACCCAGGAGTTTGGAA CAATATATTATTA GTTTAATGGTTTA TTAAATATTTTAACTTTAA GTTCAAAAC
 620 630 640 650 660 670 680 690 700 7



FIG. 2B2

APPROVED	O.G. FIG.	
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TAAA AACAA TA TTGAAAA ANTAA ATTGAA TGGG TTA CA TTTTATGAAAAAACA TCC TG ATA GCCCTT CAA ACTTCTTT CCTTTAAG
 10 720 730 740 750 760 770 780 790



CAT TTGG GOC TACA TCTCC TTA ACTTTTCTANCOONIECCTA CCTT TGGC TNACTGGATTTTTIN
 800 810 820 830 840 850 860



FIG. 2B3